

**Selective homocysteine-lowering gene transfer attenuates pressure overload-
induced cardiomyopathy via reduced oxidative stress**

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ABSTRACT

Plasma homocysteine levels predict heart failure incidence in prospective epidemiological studies. We evaluated whether selective homocysteine lowering gene transfer beneficially affects cardiac remodelling and function in a model of pressure overload-induced cardiomyopathy induced by transverse aortic constriction (TAC). Female C57BL/6 *low-density lipoprotein receptor* (*Ldlr*^{-/-}) *cystathionine-β-synthase* (*Cbs*^{+/-}) mice were fed standard chow (control mice) or a folate-depleted, methionine-enriched diet to induce hyperhomocysteinemia (diet mice). Three weeks after initiation of this diet, mice were intravenously injected with 5×10^{10} viral particles of an E1E3E4-deleted hepatocyte-specific adenoviral vector expressing *Cbs* (AdCBS), with the same dose of control vector, or with saline buffer. TAC or sham operation was performed two weeks later. AdCBS gene transfer resulted in 86.4% ($p < 0.001$) and 84.6% ($p < 0.001$) lower homocysteine levels in diet sham mice and diet TAC mice, respectively. Mortality rate was significantly reduced in diet AdCBS TAC mice compared to diet TAC mice during a follow-up period of 8 weeks (hazard ratio for mortality 0.495, 95% CI 0.249 to 0.985). Left ventricular hypertrophy ($p < 0.01$) and interstitial myocardial fibrosis ($p < 0.001$) were strikingly lower in control TAC mice and diet AdCBS TAC mice compared to diet TAC mice. Diastolic function in diet AdCBS TAC mice was similar to that of control TAC mice and was significantly improved compared to diet TAC mice. AdCBS gene transfer potentially reduced oxidative stress as evidenced by a reduction of plasma TBARS and a reduction of myocardial 3-nitrotyrosine-positive area (%). In conclusion, selective homocysteine lowering potentially attenuates pressure overload-induced cardiomyopathy via reduced oxidative stress.

KEYWORDS

Heart failure, homocysteine, cardiomyopathy, transverse aortic constriction, gene transfer.

INTRODUCTION

Clinical data support the hypothesis that plasma homocysteine levels modulate myocardial biology and function. Plasma homocysteine levels are directly related to left ventricular mass and wall thickness in women but not in men in the Framingham Heart Study participants that were free of heart failure and previous myocardial infarction[1]. Elevated homocysteine levels are associated with reduced regional left ventricular systolic function determined by tagged magnetic resonance imaging in asymptomatic subjects[2]. Furthermore, left ventricular diastolic function is significantly worse in hypertensive patients with elevated levels of homocysteine compared to hypertensive patients with normal homocysteine levels[3]. Finally, plasma homocysteine levels predict heart failure incidence in patients with or without prior myocardial infarction[4-6]. Notwithstanding the consistency of the results of these clinical studies, a causal role of homocysteine in myocardial biology and function cannot be proven since residual confounding in multivariable models may occur.

The potential role of homocysteine on cardiac function is supported by *in vitro* and *ex vivo* experiments. In isolated cultured adult rat ventricular myocytes, exposure to homocysteine increases production of reactive oxygen species, impairs cardiomyocyte contractility in a concentration-dependent manner, and promotes apoptosis[7]. Cardiomyocytes isolated from hyperhomocysteinemic mice are characterized by impaired *ex vivo* contraction and relaxation[8]. Mild hyperhomocysteinemia induced in rats via chronic methionine administration or in heterozygous *cystathionine- β -synthase* (*Cbs*) deficient mice induces cardiomyocyte hypertrophy and interstitial fibrosis[9,10]. However, an analysis of *in vivo* cardiac function has not been performed in these models. Recently, we demonstrated that selective homocysteine lowering gene transfer in female C57BL/6 *low-density lipoprotein receptor* (*Ldlr*)^{-/-} *Cbs*^{+/-} mice fed a folate-depleted, methionine-enriched diet improves infarct healing, attenuates adverse ventricular remodelling, and significantly enhances diastolic function following permanent ligation of the left

1 anterior descending coronary artery[11]. However, since beneficial effects of selective
2 homocysteine lowering gene therapy on cardiac function post-ligation of the left anterior
3 descending coronary artery may be entirely dependent on improved infarct healing and reduced
4 infarct expansion, direct effects of homocysteine levels on the ventricular wall cannot be proven
5 in this model of myocardial infarction.
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10 Transverse aortic constriction (TAC) is a commonly used model of pressure overload-induced
11 cardiac hypertrophy and heart failure[12]. TAC initially leads to compensatory hypertrophy of the
12 heart, but over time, the response to chronic hemodynamic overload becomes maladaptive and
13 results in cardiac dilatation and heart failure with reduced ejection fraction. The objective of the
14 current study is to evaluate whether plasma homocysteine has direct effects on the myocardium
15 by investigating the impact of selective homocysteine lowering gene transfer on the development
16 of pressure overload-induced cardiomyopathy in mice. Specifically, we investigated the
17 hypothesis that lower homocysteine levels inhibit the development of non-ischemic
18 cardiomyopathy via reduced oxidative stress.
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MATERIALS AND METHODS

For detailed methodology, please see the Supplementary Materials and Methods in the Electronic Supplementary Material.

Construction, generation, and production of E1E3E4-deleted adenoviral gene transfer vectors-

The construction of the E1E3E4-deleted adenoviral vector AdCBS, which induces hepatocyte-specific expression of cystathionine- β -synthase (CBS), has been described previously[13]. This vector contains the 1.27 kb DC172 promoter[14], consisting of an 890 bp human α_1 -antitrypsin promoter and two copies of the 160 bp α_1 -microglobulin enhancer, upstream of the 5' untranslated region of the human *apo A-I* gene that contains the first intron, the 1.7 kb cDNA sequence of murine *Cbs*, and 2 copies of the 774 bp *hepatic control region-1*. Hepatic overexpression of murine CBS after AdCBS gene transfer in C57BL/6 *Ldlr*^{-/-} *Cbs*^{+/-} mice has been previously confirmed by Western blot[11]. The E1E3E4-deleted control vector Adnull does not contain an expression cassette[15]. Large scale vector production was performed as described previously[15].

In vivo experiments-All experimental procedures in animals were performed in accordance with protocols approved by the Institutional Animal Care and Research Advisory Committee of the Catholic University of Leuven (Approval number: P154/2013). Control female C57BL/6 *Ldlr*^{-/-} *Cbs*^{+/-} mice[13] were fed standard chow (Sniff Spezialdiäten GMBH, Soest, Germany). Diet mice received a folate-depleted, methionine-enriched diet (TD00205; 0.2 mg/kg folic acid, 4.1 g/kg L-methionine; Harlan Teklad, Horst, The Netherlands) starting from the age of 12 weeks to induce hyperhomocysteinemia. Three weeks after initiation of this diet, tail vein injection was performed with 5×10^{10} adenoviral particles of AdCBS, with the same dose of the control vector Adnull, or with saline buffer. The equivalency of Adnull and saline injection with regard to different outcome measures has been demonstrated in several studies[16-18,11]. The experimental diet was

maintained throughout the entire duration of the experiments. Taken together, control mice were compared with diet mice and diet AdCBS mice.

To induce pressure overload, transverse aortic constriction (TAC) was performed two weeks after gene transfer or saline injection as previously described[12,19]. The sham procedure was identical except that no constriction on the aorta was applied.

Biochemical endpoint analyses-A detailed description of these quantifications is included in the Supplementary Materials and Methods.

In vivo hemodynamic measurements-Invasive hemodynamic measurements were performed 8 weeks after TAC or after sham operation as described before[17]. A detailed description can be found in the Supplementary Materials and Methods.

Histological and morphometric analysis-After hemodynamic analysis, mice were perfused via the abdominal aorta with phosphate-buffered saline (PBS) and hearts were arrested in diastole by CdCl₂ (100 μ l; 0.1 N), followed by perfusion fixation with 1% paraformaldehyde in phosphate buffered saline. After dissection, hearts were post-fixated overnight in 1% paraformaldehyde, embedded in paraffin, and 6 μ m thick cross-sections at 130 μ m spaced intervals were made extending from the apex to the basal part of the left ventricle. Left ventricle (LV) remodelling was assessed by morphometric analysis on mosaic images of Sirius red-stained heart cross-sections using Axiovision 4.6 software (Zeiss, Zaventem, Belgium). Whole LV area (mm²), LV cavity area (mm²), LV remote muscle area including the entire septum (mm²), anterior wall thickness, and septal wall thickness were analyzed. All geometric measurements were computed in a blinded fashion from representative tissue sections of 4 separate regions and the average value was used to represent that animal for statistical purposes.

A detailed description of (immuno)histochemical analyses is included in the Supplementary Materials and Methods.

Statistical analysis-All data are expressed as means \pm standard error of the means (SEM).

Detailed description of statistical analysis can be retrieved in the Supplementary Materials and Methods.

RESULTS

AdCBS gene transfer selectively lowers homocysteine levels in diet mice

Plasma homocysteine levels were 15.5-fold ($p<0.001$) and 19.4-fold ($p<0.001$) higher in diet sham and diet TAC C57BL/6 *Ldlr*^{-/-} *Cbs*^{+/-} mice compared to respective control groups (Figure 1). AdCBS gene transfer resulted in 86.4% ($p<0.001$) and 84.6% ($p<0.001$) lower homocysteine levels in diet sham mice and diet TAC mice, respectively. No significant difference of plasma homocysteine levels was observed between control groups and diet AdCBS groups (Figure 1). As illustrated in Supplementary Table 1, there was no impact of the diet, of AdCBS gene transfer, or of the TAC procedure on plasma lipoprotein cholesterol levels.

Selective homocysteine lowering gene transfer improves survival after TAC in diet mice

Comparison of Kaplan-Meier survival curves (Figure 2) showed a higher mortality rate in diet TAC mice compared to TAC control mice ($p=0.0519$) (hazard ratio for mortality 1.97, 95% CI 0.994 to 3.89). Mortality rate was significantly reduced in diet AdCBS TAC mice compared to diet TAC mice ($p<0.05$) (hazard ratio for mortality 0.495, 95% CI 0.249 to 0.985) (Figure 2). No significant difference in mortality rate was observed between control TAC mice and diet AdCBS TAC mice (Figure 2). Sham operation did not result in any mortality (data not shown).

Cardiac hypertrophy is lower and interstitial fibrosis is reduced in diet AdCBS TAC mice compared to diet TAC mice

No significant difference in heart weights (Figure 3A) or lung weights (Figure 3B) was observed between the different sham groups. The heart weight was 2.55-fold ($p<0.0001$) higher in diet TAC mice compared to diet sham mice (Figure 3A). Compared to diet TAC mice, cardiac hypertrophy was 25.3% ($p<0.01$) and 16.0% ($p<0.05$) lower in control TAC mice and diet AdCBS TAC mice, respectively. The lung weight was 2.07-fold ($p<0.001$) higher in diet TAC mice compared to diet sham mice (Figure 3B). Lung weight was 32.8% ($p<0.05$) and 15.5%

(p=NS) lower in control TAC mice and diet AdCBS TAC mice, respectively, than in diet TAC mice.

Morphometric and histological parameters of the left ventricular myocardium 8 weeks after sham operation are shown in Table 1. No significant differences were observed between the three groups with the exception of a slight increase of interstitial fibrosis in diet sham mice and diet AdCBS sham mice compared to sham control mice. Morphometric and histological parameters of the left ventricular myocardium 8 weeks after TAC are illustrated in Table 2. In agreement with the heart weight data (Figure 3A), LV wall area, septal wall thickness, and anterior wall thickness were significantly lower in control TAC mice and diet AdCBS TAC mice compared to diet TAC mice (Table 2, Supplementary Figure 1). Cardiomyocyte hypertrophy was significantly more pronounced in the diet TAC mice compared to the two other groups (Table 2, Supplementary Figure 2). Capillary density and relative vascularity were 22.9% ($p<0.05$) and 24.4% ($p=NS$) higher, respectively, in diet AdCBS TAC mice compared to diet TAC mice (Table 2, Supplementary Figure 2). Interstitial fibrosis was strikingly lower in control TAC mice and diet AdCBS TAC mice compared to diet TAC mice (Table 2, Supplementary Figure 2). Apoptosis in the myocardium was evaluated using immunohistochemical quantification of cleaved caspase-3. Cleaved caspase-3 positive cells were undetectable in the myocardium of control sham mice and diet AdCBS sham mice whereas cleaved caspase-3 positive cells were infrequently observed in diet sham mice (Table 1). Compared to diet TAC mice, the number of cleaved caspase-3 positive cells was reduced by 36.0% ($p<0.01$) and 32.9% ($p<0.01$) in control TAC mice and diet AdCBS TAC mice, respectively (Table 2).

Selective homocysteine lowering gene transfer improves diastolic function after TAC

Hemodynamic parameters in the left ventricle and in the aorta 8 weeks after sham operation are summarized in Table 3. There was a 15.7% ($p<0.01$) decrease of the absolute value of the peak rate of isovolumetric relaxation (dP/dt_{min}) in diet sham mice compared to control sham mice.

Following TAC, diastolic function in diet AdCBS mice was similar compared to control mice and was significantly improved compared to diet mice as evidenced by the higher absolute value of dP/dt_{min} and a lower value of the time constant of left ventricular relaxation (Table 4). End-diastolic pressure was significantly lower in control TAC mice and diet AdCBS TAC mice than in diet TAC mice (Table 4).

Hyperhomocysteinemia reduces antioxidants and decreases antioxidant defense systems in plasma

To quantify antioxidants and antioxidant defense systems, the concentration of glutathione (GSH) and cysteine and the activity of glutathione peroxidase and superoxide dismutase were determined in plasma (Supplementary Table 2 and Supplementary Table 3). The concentration of glutathione (GSH) and cysteine were significantly reduced in diet sham mice compared to control sham mice and diet sham AdCBS mice (Supplementary Table 2). Furthermore, glutathione peroxidase activity and superoxide dismutase activity were significantly lower in diet sham mice compared to the other sham groups (Supplementary Table 2). As shown in Supplementary Table 3, antioxidants and antioxidant defense systems were significantly decreased in diet TAC mice compared to control TAC mice and diet AdCBS TAC mice.

AdCBS gene transfer reduces oxidative stress

Reduction of oxidative stress may underlie the beneficial effects of selective homocysteine lowering gene transfer on cardiac hypertrophy, interstitial fibrosis, and diastolic function. Compared to diet sham mice, plasma TBARS were 32.5% ($p<0.05$) lower in both control sham mice and diet AdCBS sham mice (Figure 4A). TBARS were increased by 30.3% ($p=0.0542$), 126% ($p<0.01$), and 63.4% ($p<0.01$) in control TAC mice, diet TAC mice, and diet AdCBS TAC mice, respectively, compared to respective sham groups. Furthermore, TBARS were significantly

($p < 0.001$) higher in the diet TAC mice compared to the two other TAC groups (Figure 4A). Plasma asymmetric dimethylarginine (ADMA) levels were 26.6% ($p < 0.05$) and 27.4% ($p < 0.05$) higher in control sham mice and in diet AdCBS sham mice, respectively, than in diet sham mice (Figure 4B). Compared to respective sham groups, a significant increase of ADMA occurred in all TAC groups but no significant difference was observed between TAC groups (Figure 4B). Representative myocardial sections immunostained for 3-nitrotyrosine staining are shown in Figure 4C. Compared to respective sham groups, 3-nitrotyrosine-positive area (%) was increased 4.65-fold ($p < 0.0001$), 3.78-fold ($p < 0.001$), and 2.69-fold ($p < 0.0001$) in control TAC mice, diet TAC mice, and diet AdCBS TAC mice, respectively (Figure 4D). The 3-nitrotyrosine-positive area was 54.7% ($p < 0.01$) and 38.6% ($p < 0.05$) lower in control TAC mice and in diet AdCBS TAC mice, respectively, than in diet TAC mice. Taken together, selective homocysteine lowering gene transfer reduces myocardial oxidative stress.

DISCUSSION

The main findings of the current study are that 1) selective homocysteine lowering gene therapy in a murine model of hyperhomocysteinemia significantly reduces mortality after TAC; 2) lower plasma homocysteine levels result in reduced cardiac hypertrophy, decreased interstitial fibrosis, and improved diastolic function after TAC compared to hyperhomocysteinemic mice; 3) lower homocysteine levels result in significantly improved antioxidant defense systems both in sham mice and in TAC mice; 4) lower homocysteine plasma levels reduce oxidative stress in both sham mice and in TAC mice as evidenced by a reduction of plasma TBARS and myocardial 3-nitrotyrosine levels; 5) apoptosis in the myocardium is reduced in control TAC mice and diet AdCBS TAC mice compared to diet TAC mice.

Most studies on the effect of homocysteine on biochemical, biological, and medical end-points depend on dietary manipulations that may have effects independent of homocysteine levels. An important strength of the current study is that the decrease of homocysteine levels in mice on the folate-depleted, methionine-enriched diet was induced by selective homocysteine lowering gene transfer. Therefore, dietary effects unrelated to homocysteine lowering cannot have an impact on the end-points in our study. Most but not all parameters were very similar in control TAC mice and in diet AdCBS TAC mice indicating that in general the impact of hyperhomocysteinemia overrides any effect resulting from homocysteine-unrelated differences between standard chow and the folate-depleted, methionine-enriched diet.

Lower homocysteine levels were associated with lower lung weights at 8 weeks after TAC, which is indicative of decreased congestive heart failure. Furthermore, mortality in diet AdCBS TAC mice was significantly reduced compared to diet TAC mice, which predominantly reflects enhanced survival in the first four weeks after TAC. Early mortality after TAC may be caused by acute heart failure or by lethal arrhythmias[20].

Lower plasma homocysteine levels resulted in reduced cardiac hypertrophy after TAC. According to the classic wall stress-hypothesis of Grosmann[21], pressure overload induces myocytes to grow in width until wall thickness has increased sufficiently to normalize systolic wall stress. In this concentric hypertrophy, the fibers get thicker because sarcomeres are laid down in parallel[22]. However, hypertrophy is also influenced by load-independent factors as demonstrated by the effect of genetic background[23] and as illustrated by observations in many transgenic models[24]. Notably, these studies[23,24] highlight that there is no direct relationship between the degree of cardiac hypertrophy and cardiac function. In a mouse model with transgenic inactivation of G_q , hypertrophy from pressure loading was less than predicted and wall stress was not normalized[25]. Contractility in these transgenic mice was better than in wild-type mice with more hypertrophy and normalization of wall stress. On the other hand, dual specificity mitogen-activated protein kinase kinase 1 (MEK1) transgenic mice demonstrated concentric hypertrophy without signs of cardiomyopathy or lethality up to 12 months of age[26]. These MEK1 transgenic mice showed a dramatic increase in cardiac function as quantified by echocardiography and as measured in isolated working heart preparations[26]. Here, we demonstrate that reduced cardiac hypertrophy following lowering of plasma homocysteine levels is associated with improved diastolic function and a trend for an improvement of systolic function. The relation between cardiac hypertrophy and cardiac function can also be analyzed from a clinical perspective. Left ventricular hypertrophy has an important prognostic value in patients. Hypertensive patients with concentric left ventricular hypertrophy have the highest incidence of cardiovascular events including death[27]. In patients with severe aortic stenosis, the degree of left ventricular hypertrophy independently predicts heart failure regardless of the degree of flow restriction imposed by the valve pathology[28]. Taken together, less pronounced left ventricular hypertrophy as observed following homocysteine lowering may be beneficial and may contribute to the improved cardiac function. In this regard, homocysteine levels in patients correlate with

1 brain natriuretic peptide[29] and the N-terminal of the prohormone brain natriuretic peptide[30],
2 biomarkers of chronic heart failure.
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4 Pressure overload resulted in oxidative stress in all three TAC groups as evidenced by elevated
5 plasma TBARS and 3-nitrotyrosine immunostaining. Oxidative stress refers to situations
6 characterized by an imbalance between the production of reactive oxygen species and antioxidant
7 defenses leading to oxidative modification of numerous biological substrates[31]. A chronic
8 increase in reactive oxygen species plays a critical role in the development and progression of
9 heart failure. Markers of oxidative stress are increased in patients with heart failure and correlate
10 with the severity of cardiac dysfunction[32]. Reactive oxygen species can induce DNA damage,
11 protein damage, and disrupt cell membranes. They may contribute to cardiomyocyte hypertrophy
12 and lead to cardiomyocyte apoptosis. Reactive oxygen species can be produced through electron
13 leakage from the mitochondria during oxidative phosphorylation. Alternatively, they are
14 generated by several enzymes including nicotinamide adenine dinucleotide phosphate (NADPH)-
15 oxidase, xanthine oxidase, and nitric oxide synthase[33].
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17 Hyperhomocysteinemia augmented oxidative stress in TAC mice. Antioxidants (glutathione,
18 cysteine) and antioxidant defense systems (glutathione peroxidase, superoxide dismutase) were
19 significantly lower in hyperhomocysteinemic mice. CBS converts homocysteine to cystathionine.
20 The increase of cysteine and glutathione following AdCBS gene transfer reflects increased
21 substrate availability of metabolic pathways downstream of cystathionine. Handy *et al.*[34]
22 previously demonstrated that homocysteine decreases glutathione peroxidase activity by altering
23 the specific translational mechanism essential for the synthesis of this selenocysteine-containing
24 protein. Superoxide dismutase protein levels in the heart were reduced in a dog model[35] and in
25 a rat model of chronic hyperhomocysteinemia[36]. Production of reactive oxygen species is
26 increased by hyperhomocysteinemia. Auto-oxidation of homocysteine generates reactive oxygen
27 species including superoxide, hydrogen peroxide, and hydroxyl radicals, and also directly
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1 contributes to the depletion of nitric oxide via formation of peroxynitrite (ONOO⁻)[37,38]. In
2 addition, hyperhomocysteinemia has been shown to induce NADPH-oxidase 2 (Nox2) [35] and
3 Nox 4 in cardiomyocytes[39]. Furthermore, hyperhomocysteinemia promotes expression of
4 inducible NO synthase (iNOS), further increasing superoxide radical formation[40]. In isolated
5 cultured adult rat ventricular myocytes, exposure to homocysteine for 24 hours impaired
6 cardiomyocyte contractility in a concentration-dependent manner and promoted apoptosis[7].
7 These effects were associated with activation of p38 mitogen-activated protein kinase and
8 increased production of reactive oxygen species. Inhibition of p38 mitogen-activated protein
9 kinase prevented all of the effects of homocysteine[7]. Furthermore, oxidative stress activates
10 Ca²⁺-calmodulin kinase II. Under physiological conditions, this enzyme couples increases in
11 intracellular Ca²⁺ to activation of ion channels by phosphorylation of a large and diverse set of
12 effector targets[41]. As a result of oxidative stress, Ca²⁺-calmodulin kinase II activity is sustained,
13 independent of Ca²⁺ and calmodulin amounts, which may induce cardiac arrhythmias[42]. This
14 may contribute to increased mortality in diet TAC mice. This enzyme also provides a link
15 between increased oxidative stress and increased apoptosis in diet TAC mice since knockdown of
16 Ca²⁺-calmodulin kinase II in cardiac myocytes reduces apoptosis[42]. Oxidative stress may also
17 contribute to increased interstitial fibrosis in diet TAC mice. Exposure of cardiac fibroblasts to
18 superoxide anion enhances the production of transforming growth factor-β1 (TGF-β1)[43,44],
19 which is a potent fibrogenic cytokine. Recently, up-regulation of Nox4 in the myocardium has
20 been directly linked to increased interstitial fibrosis[45].

21 Asymmetric dimethylarginine (ADMA) is an endogenous competitive inhibitor of nitric oxide
22 (NO) synthase (NOS). ADMA is produced from the proteolysis of proteins that contain
23 dimethylated arginine residues. ADMA levels were quite surprisingly lower in diet sham mice
24 compared to the two sham groups with lower homocysteine levels. This observation may be the
25 result of a large decrease in the hepatic methylation capacity (low S-adenosylmethionine-to-S-

adenosylhomocysteine ratio) induced by the high-methionine low-folate diet in *Cbs*^{+/-} mice [46].

ADMA levels were increased in all three TAC groups compared to respective sham groups. These increments reflect the increased oxidative stress induced by TAC. Indeed, increased oxidative stress results in downregulation of dimethylarginine dimethylaminohydrolase[47-49]. Since dimethylarginine dimethylaminohydrolase hydrolyzes ADMA to citrulline and dimethylamine, reduced activity of this enzyme results in increased ADMA levels. However, in light of similar ADMA levels in the three TAC groups, our data do not indicate that ADMA levels contribute significantly to the more pronounced deterioration of cardiac function in diet TAC mice compared to the two TAC groups with low homocysteine levels.

In conclusion, selective homocysteine lowering gene therapy in a murine model of hyperhomocysteinemia significantly reduces mortality, attenuates cardiac hypertrophy, decreases interstitial fibrosis, reduces apoptosis, and enhances diastolic function after TAC. These beneficial effects are likely closely linked to a profound reduction of oxidative stress secondary to the decrease of homocysteine levels.

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DISCLOSURES

None of the authors has a conflict of interest.

ETHICAL APPROVAL

All procedures performed in studies involving animals were in accordance with the ethical standards of the institution or practice at which the studies were conducted (Institutional Animal Care and Research Advisory Committee of the Catholic University of Leuven. Approval number: P154/2013).

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LEGENDS TO THE FIGURES

Figure 1. Plasma homocysteine levels in control, diet, and diet AdCBS mice. Sham mice and TAC mice are indicated by open bars and closed bars, respectively. Plasma samples were obtained at day 56 after sham operation or TAC.

Figure 2. Kaplan-Meier survival curves after TAC. Control mice, diet mice, and diet AdCBS mice are indicated by black lines, red lines, and blue lines respectively. Survival curves were compared by log-rank test.

Figure 3. Heart weights (panel A) and lung weights (panel B) in control, diet, and diet AdCBS mice. Sham mice and TAC mice are indicated by open bars and closed bars, respectively. Data reflect wet weights at day 56 after sham operation or TAC.

Figure 4. Quantification of oxidative stress in sham mice and in TAC mice. Sham mice and TAC mice are indicated by open bars and closed bars, respectively. (A) Plasma TBARS expressed as plasma malondialdehyde equivalents. (B) Plasma asymmetric dimethylarginine levels. (C) Representative photomicrographs showing myocardial sections stained for 3-nitrotyrosine. Scale bar represents 100 μ m. (D) Percentage of 3-nitrotyrosine-positive area in the myocardium.

Figure 1
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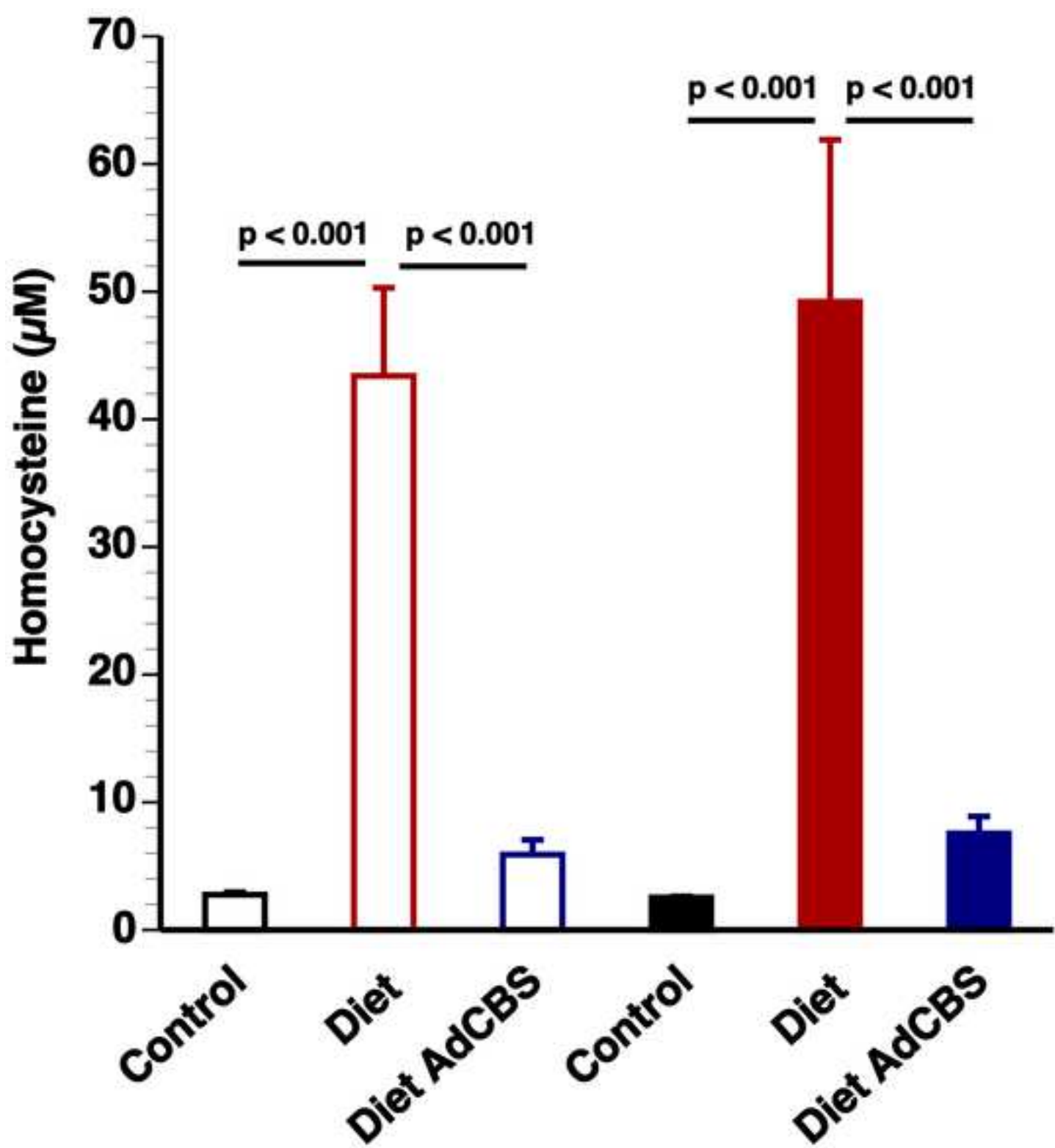


Figure 2

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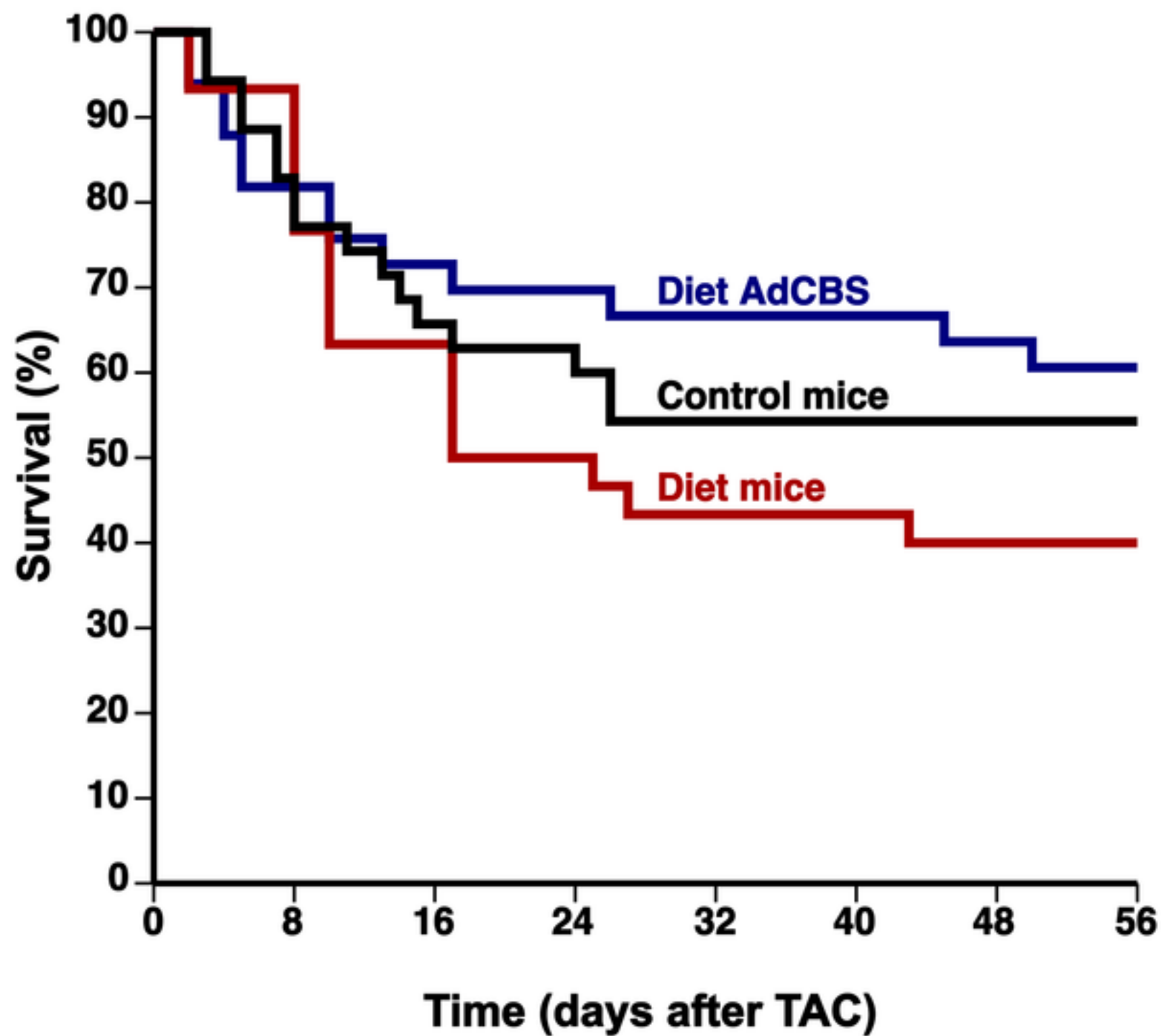


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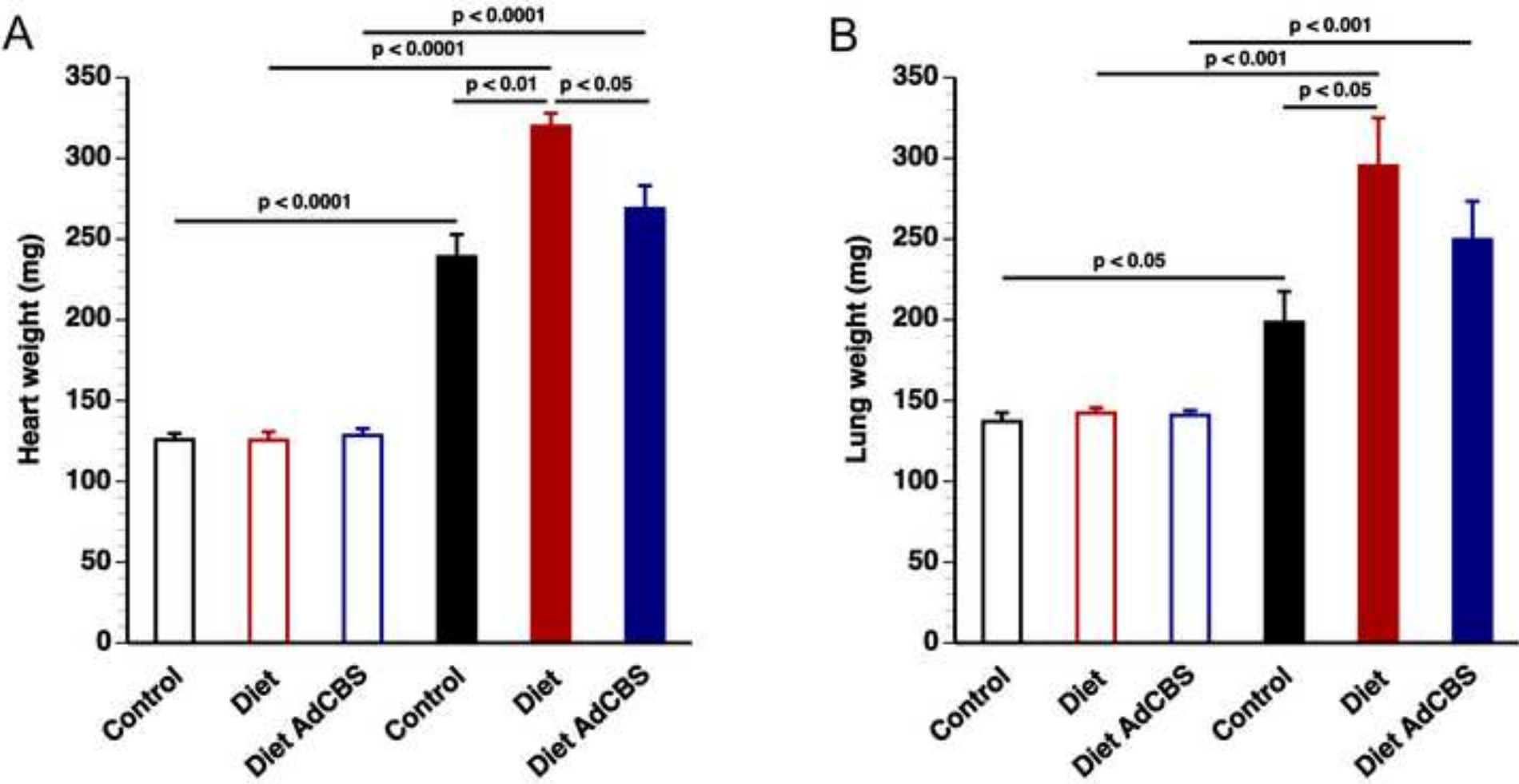
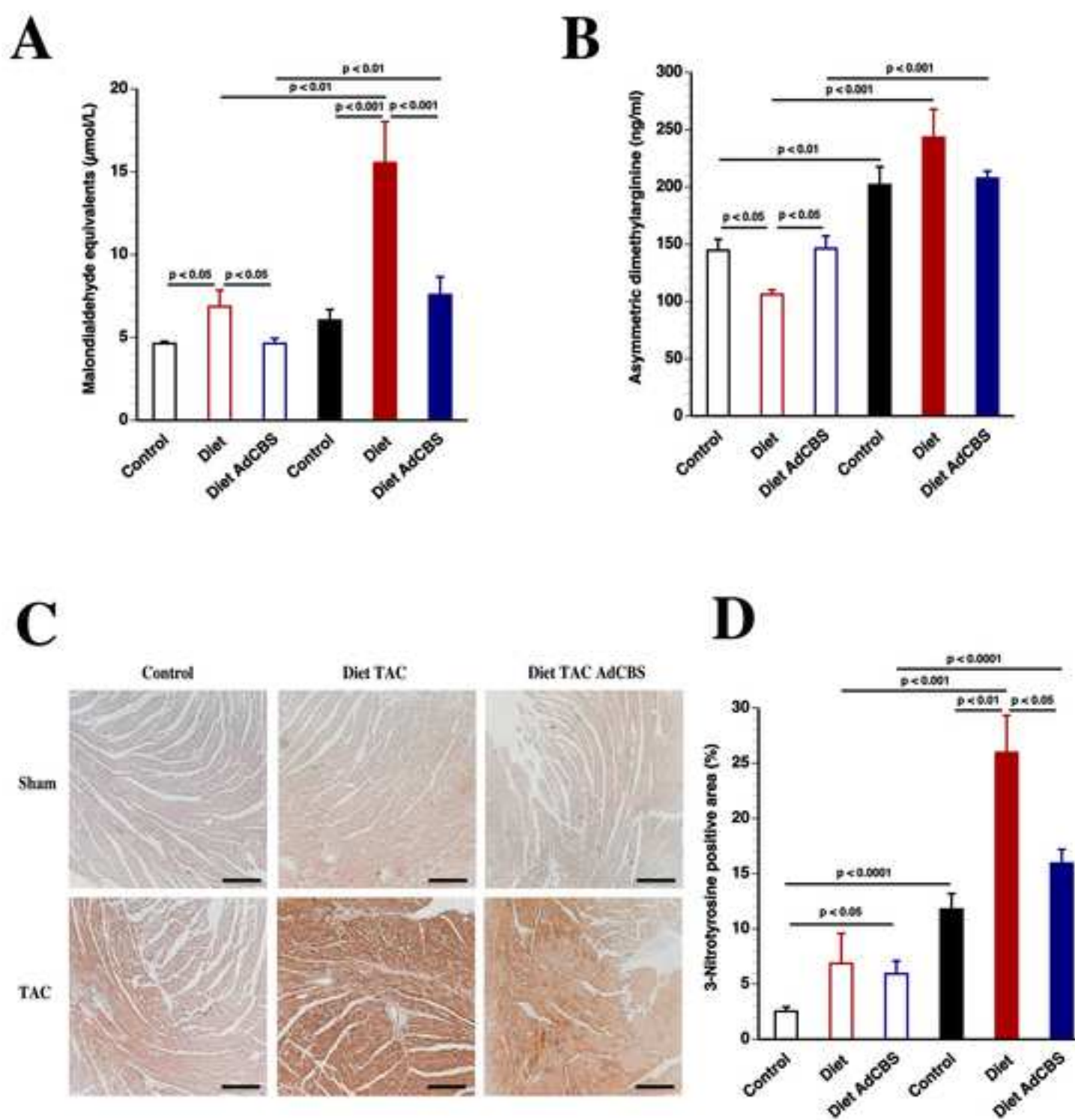
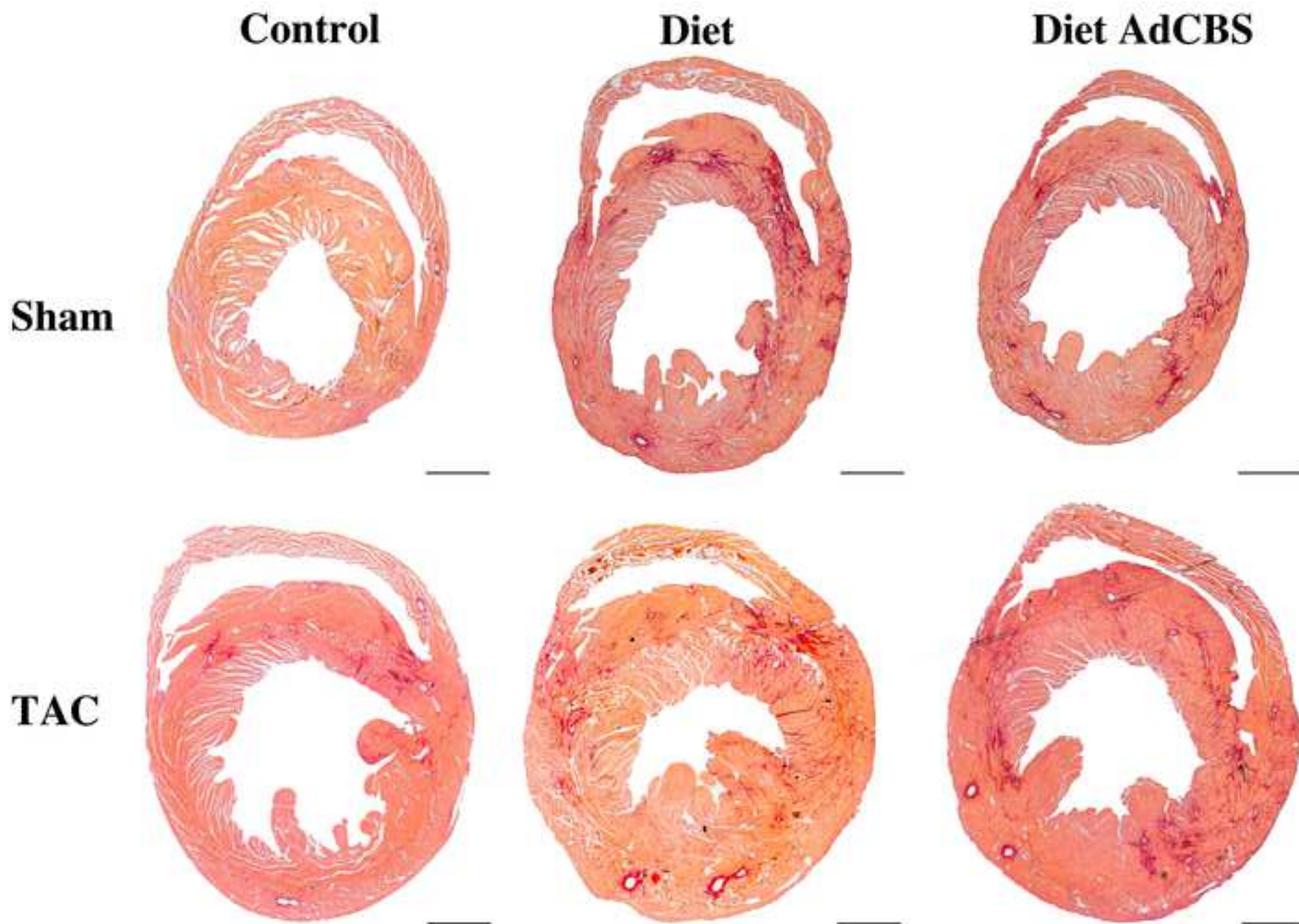


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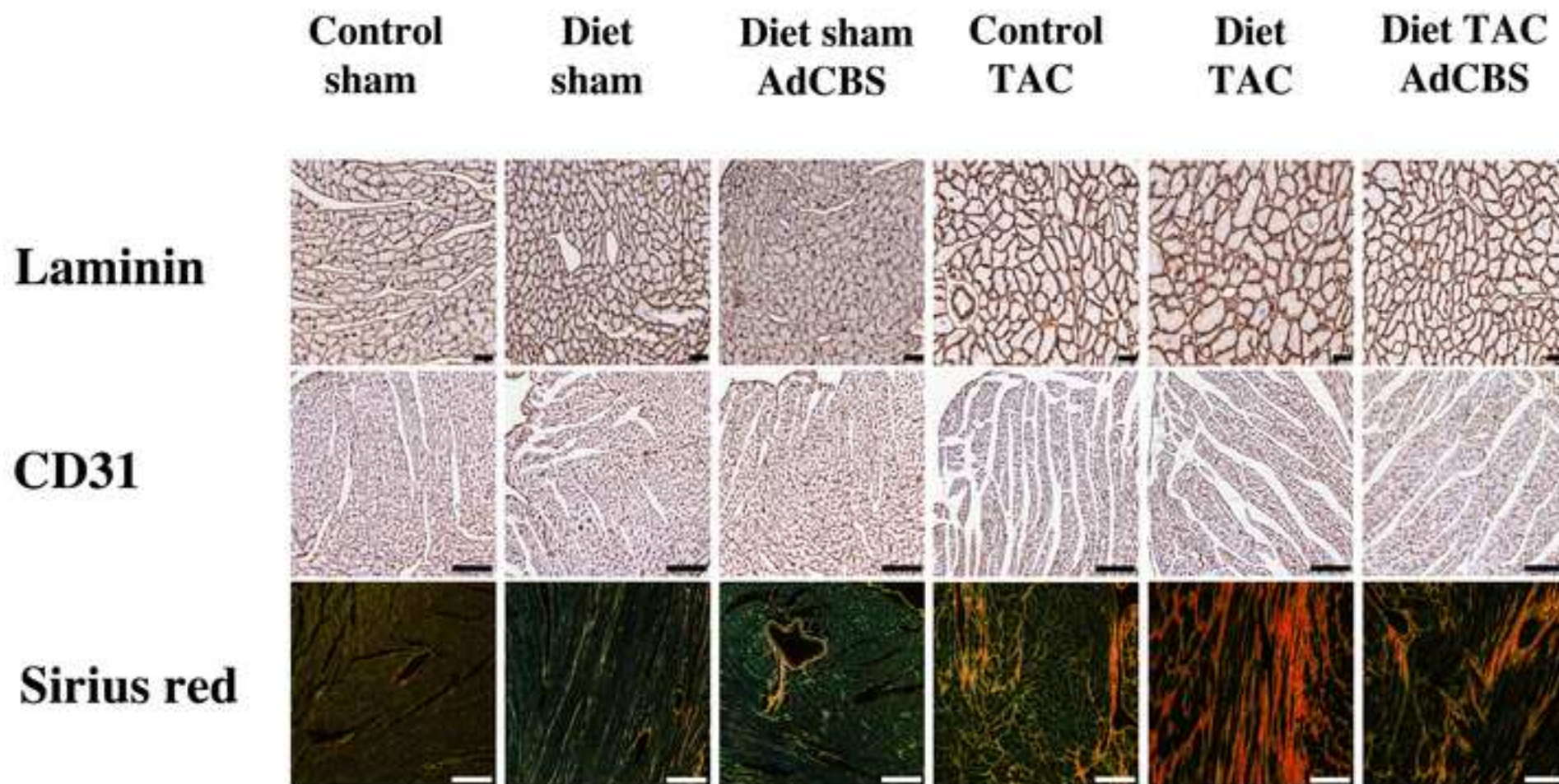


Table 1

Table 1. Morphometric and histological parameters of the left ventricular myocardium 8 weeks after sham operation in female C57BL/6 *Ldlr*^{-/-} *Cbs*^{+/-} mice.

	Control sham	Diet sham	Diet sham AdCBS
Number of mice	12	17	24
LV wall area (mm ²)	8.89 ± 0.30	9.07 ± 0.30	8.95 ± 0.11
Septal wall thickness (µm)	1100 ± 30	1110 ± 30	1090 ± 20
Anterior wall thickness (µm)	1140 ± 30	1130 ± 30	1120 ± 10
Cardiomyocyte cross-sectional area (µm ²)	215 ± 12	184 ± 8	186 ± 6
Cardiomyocyte density (number/mm ²)	5110 ± 200	5000 ± 220	5320 ± 160
Capillary density (number/mm ²)	6760 ± 250	6320 ± 170	6610 ± 150
Relative vascularity (µm ⁻²)	0.00634 ± 0.00031	0.00705 ± 0.00023	0.00684 ± 0.00020
Interstitial fibrosis (%)	2.69 ± 0.20	4.01 ± 1.3 ^{°°}	4.07 ± 0.44 [°]
Perivascular fibrosis (ratio)	0.469 ± 0.017	0.502 ± 0.026	0.489 ± 0.020
Cleaved caspase 3 positive cells (number/mm ²)	0.00 ± 0.01	2.09 ± 1.33 [°]	0.00 ± 0.01 [*]

Control mice were fed standard chow. The Diet (TD00205) is a folate-depleted, methionine-enriched diet. Data are expressed as means ± SEM. °: p<0.05; °°:p<0.01 versus Control sham. *:p<0.05 versus diet sham.

Table 2. Morphometric and histological parameters of the left ventricular myocardium 8 weeks after TAC in female C57BL/6 *Ldlr*^{-/-} *Cbs*^{+/-} mice.

	Control TAC	Diet TAC	Diet TAC AdCBS
Number of mice	17	13	23
LV wall area (mm ²)	11.1 ± 0.4	13.8 ± 0.3 ^{ooo}	12.3 ± 0.4*
Septal wall thickness (μm)	1280 ± 30	1460 ± 30 ^{ooo}	1340 ± 30**
Anterior wall thickness (μm)	1340 ± 30	1510 ± 40 ^{oo}	1320 ± 30***
Cardiomyocyte cross-sectional area (μm ²)	417 ± 25	503 ± 18 ^o	420 ± 25*
Cardiomyocyte density (number/mm ²)	3140 ± 140	2500 ± 100 ^o	3210 ± 210*
Capillary density (number/mm ²)	4560 ± 190	4360 ± 230	5360 ± 280*
Relative vascularity (μm ⁻²)	0.00372 ± 0.00024	0.00353 ± 0.00019	0.00439 ± 0.00027
Interstitial fibrosis (%)	11.3 ± 0.6	17.1 ± 1.3 ^o	9.21 ± 0.89***
Perivascular fibrosis (ratio)	0.567 ± 0.040	0.727 ± 0.042 ^o	0.529 ± 0.020**
Cleaved caspase 3 positive cells (number/mm ²)	15.8 ± 1.5	24.7 ± 2.0 ^{oo}	16.6 ± 1.0**

Control mice were fed standard chow. The diet (TD00205) is a folate-depleted, methionine-enriched diet.

Data are expressed as means ± SEM. °: p<0.05; °°:p<0.01; °°°:p<0.001 versus Control TAC. *:p<0.05; **:p<0.01; ***:p<0.001 versus Diet TAC.

Table 3. Hemodynamic parameters in the left ventricle and in the aorta 8 weeks after sham operation in female C57BL/6 *Ldlr*^{-/-} *Cbs*^{+/-} mice.

	Control sham	Diet sham	Diet sham AdCBS
Number of mice	8	17	20
LEFT VENTRICLE			
Peak systolic pressure (mm Hg)	103 ± 3	99.1 ± 1.4	99.4 ± 2.3
End-diastolic pressure (mm Hg)	3.07 ± 0.45	1.69 ± 0.49	1.69 ± 0.36
dP/dt max (mm Hg/ms)	13.6 ± 0.8	11.6 ± 0.6	11.9 ± 0.4
dP/dt min (mm Hg/ms)	-11.4 ± 0.5	-9.61 ± 0.36 ^{°°}	-10.1 ± 0.2
Tau (ms)	4.46 ± 0.19	4.83 ± 0.19	4.83 ± 0.12
Heart rate (bpm)	598 ± 18	611 ± 10	571 ± 11
AORTA			
Mean pressure (mm Hg)	97.7 ± 2.6	101 ± 5	98.7 ± 3.3
Peak systolic pressure (mm Hg)	101 ± 2	98.7 ± 1.7	99.3 ± 2.9
Peak diastolic pressure (mm Hg)	64.9 ± 3.4	69.6 ± 2.9	67.6 ± 4.0

Control mice were fed standard chow. The diet (TD00205) is a folate-depleted, methionine-enriched diet. Hemodynamic analysis was performed at the age of 25 weeks. Data are expressed as means ± SEM. °°:p<0.01 versus Control sham.

Table 4. Hemodynamic parameters in the left ventricle and in the aorta 8 weeks after TAC in female C57BL/6 *Ldlr*^{-/-} *Cbs*^{+/-} mice.

	Control TAC	Diet TAC	Diet TAC AdCBS
Number of mice	14	10	12
LEFT VENTRICLE			
Peak systolic pressure (mm Hg)	162 ± 6	159 ± 8	164 ± 4
End-diastolic pressure (mm Hg)	1.88 ± 0.55	6.26 ± 0.80 ^{°°°}	3.77 ± 0.37**
dP/dt _{max} (mm Hg/ms)	9.48 ± 0.43	8.55 ± 0.71	9.55 ± 0.36
dP/dt _{min} (mm Hg/ms)	-9.25 ± 0.58	-7.78 ± 0.41 [°]	-9.02 ± 0.25*
Tau (ms)	5.72 ± 0.22	7.07 ± 0.30 ^{°°}	5.91 ± 0.24*
Heart rate (bpm)	576 ± 21	580 ± 31	585 ± 15
AORTA			
Mean pressure (mm Hg)	97.7 ± 2.6	101 ± 5	98.2 ± 3.8
Peak systolic pressure (mm Hg)	159 ± 5	157 ± 8	163 ± 5
Peak diastolic pressure (mm Hg)	59.5 ± 3.5	67.2 ± 4.7	58.6 ± 2.9

Control mice were fed standard chow. The diet (TD00205) is a folate-depleted, methionine-enriched diet. Hemodynamic analysis was performed at the age of 25 weeks. Data are expressed as means ± SEM. °:p<0.05; °°:p<0.01; °°°:p<0.001 versus Control TAC. *:p<0.05; **:p<0.01 versus Diet TAC.

Supplementary Material

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